



Fig. S5. Simplified RAxML phylogeny of 133 accessions representing 35 species of *Salix* subg. *Chamaetia/Vetrix* based on 1,660 RAD loci resulting from the m100 dataset. Ploidy level is indicated behind species name, polyploid samples are highlighted in bold. *Salix triandra* (subg. *Salix*) was used to root the tree. BS values above branches.